

FIGURE 1A

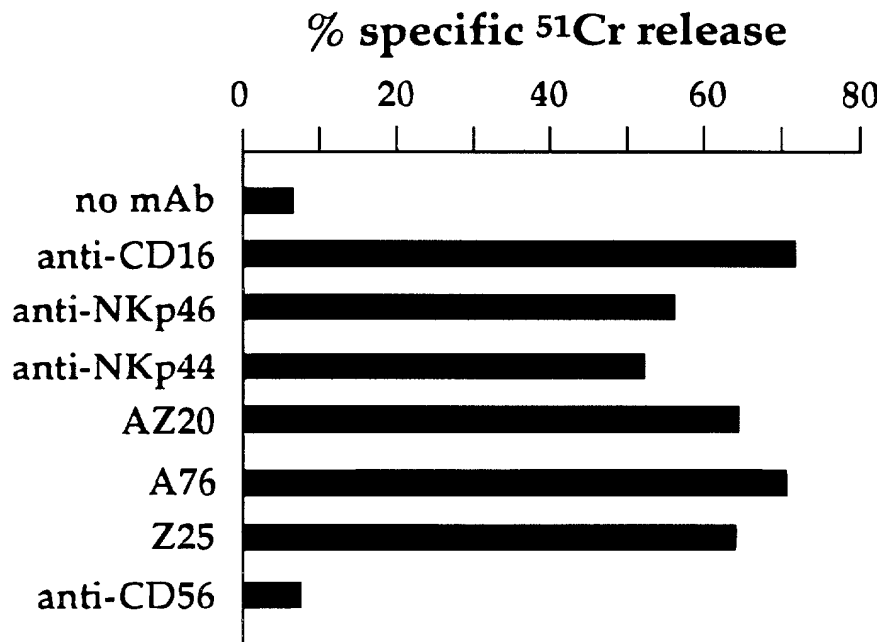


FIGURE 1B

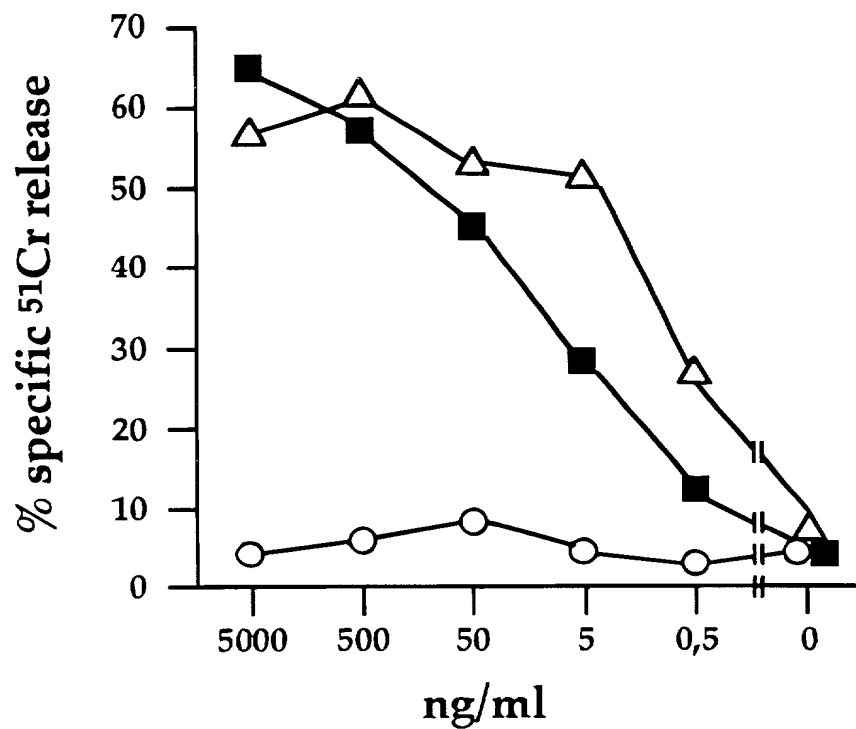


FIGURE 1C

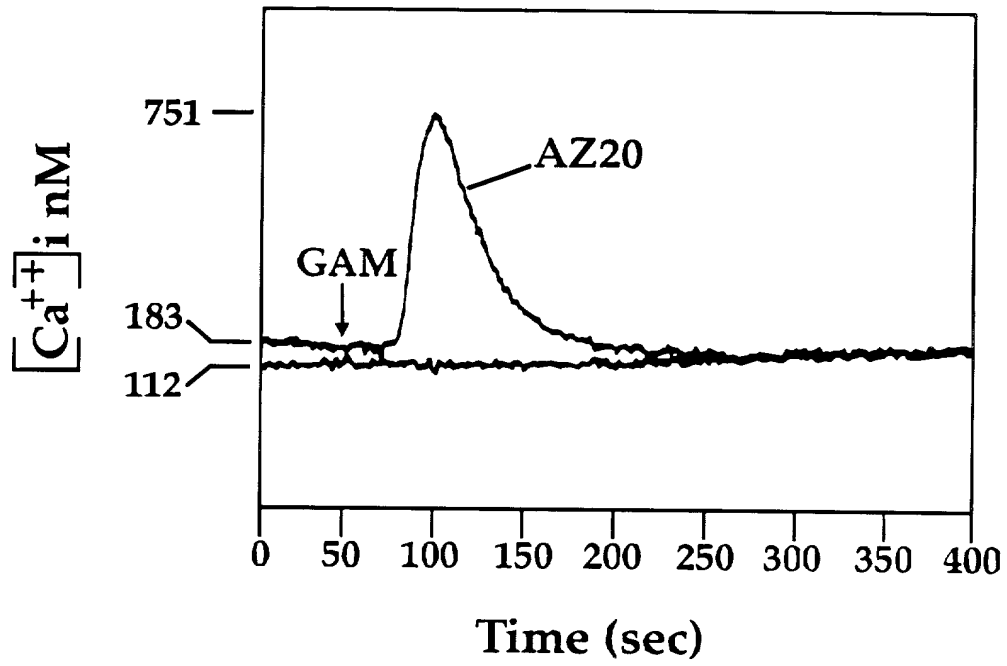


FIGURE 2A

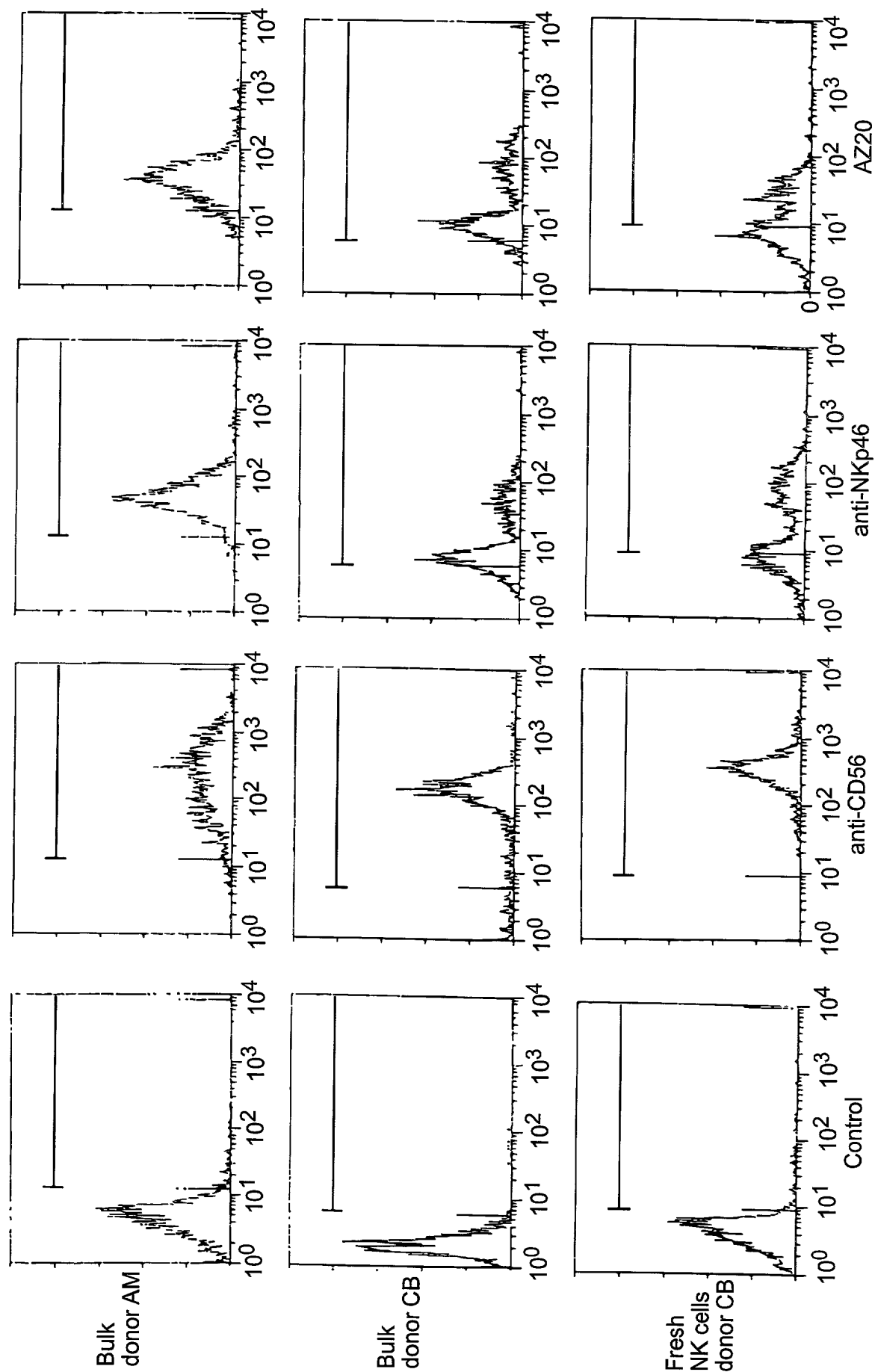


FIGURE 2B

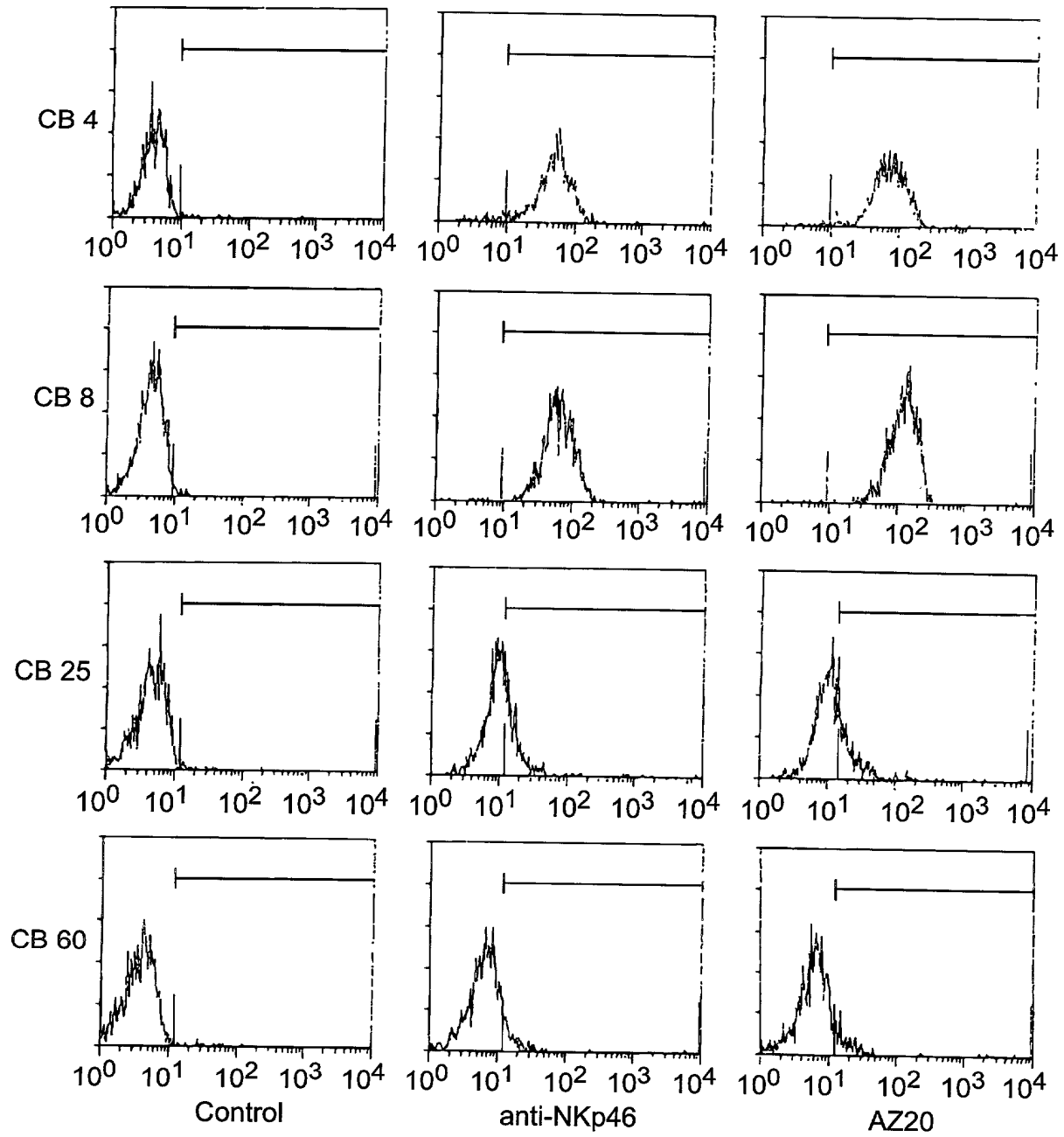


FIGURE 3A

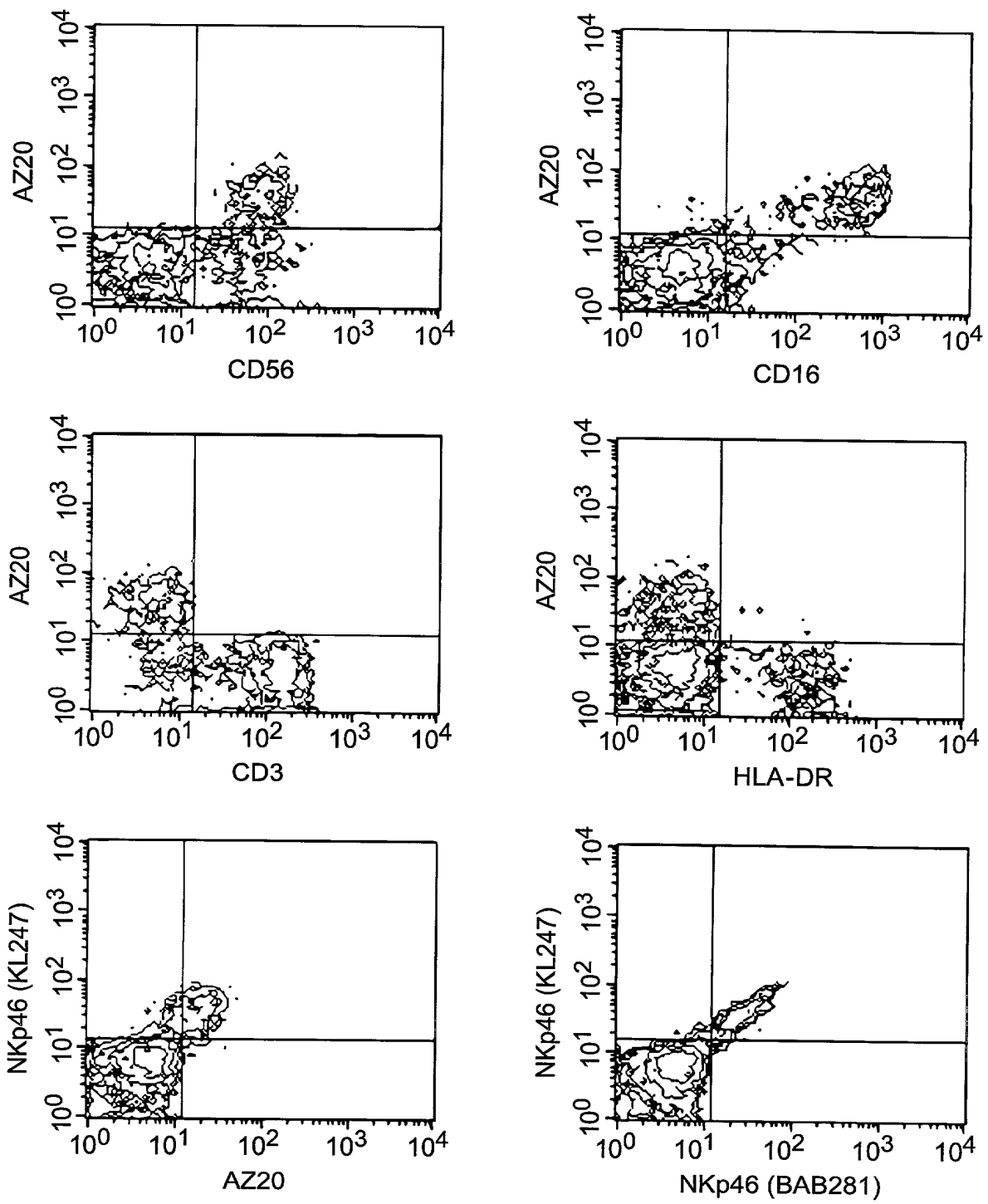


FIGURE 3B

Cells Daudi NK

69 —

45 —

28 —

18 —

Blot

AZ20



FIGURE 4A

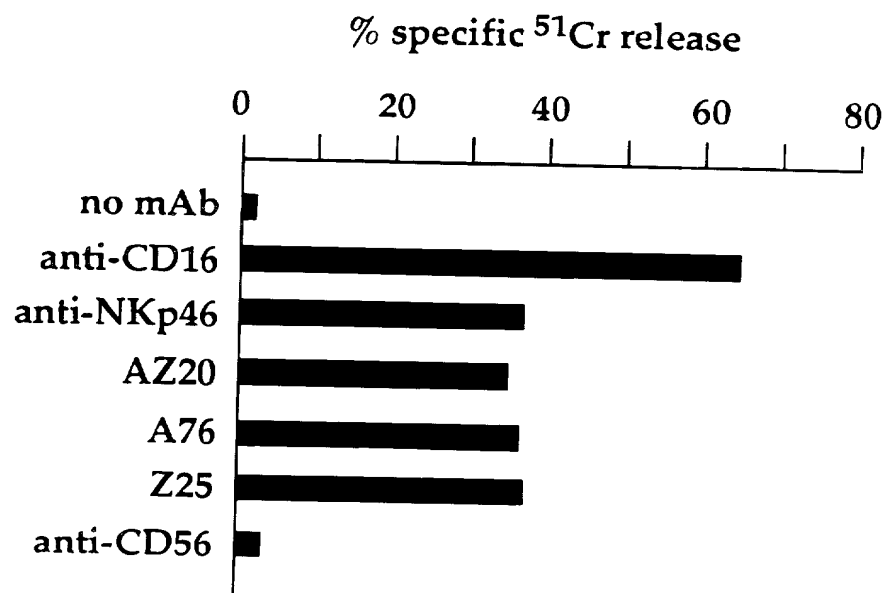


FIGURE 4B

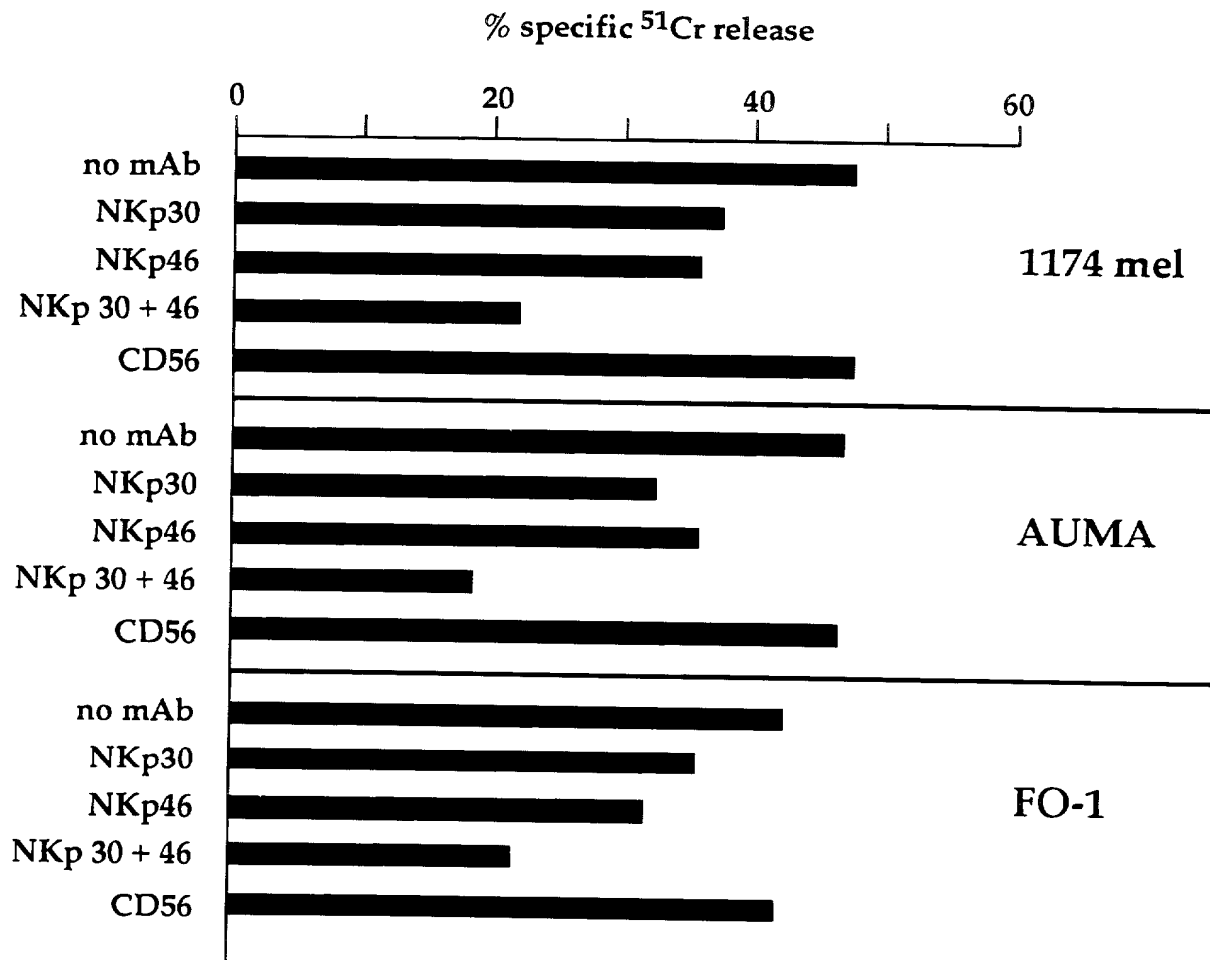


FIGURE 5

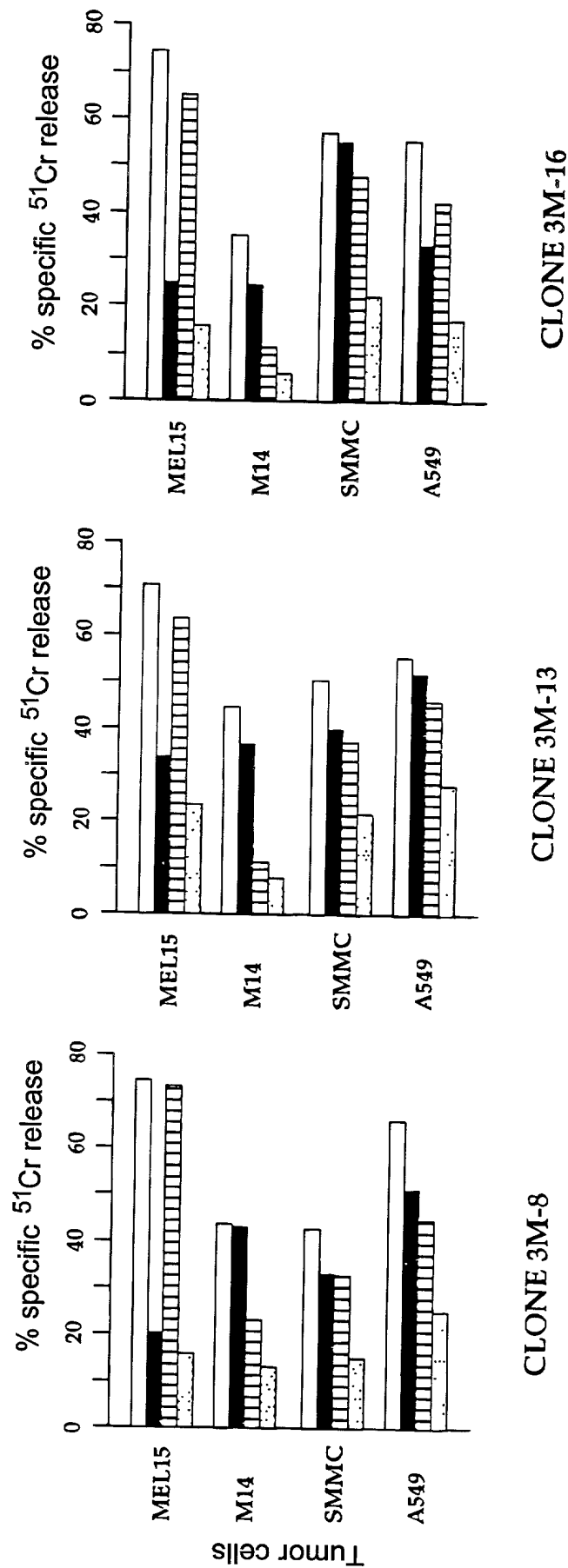


FIGURE 6A

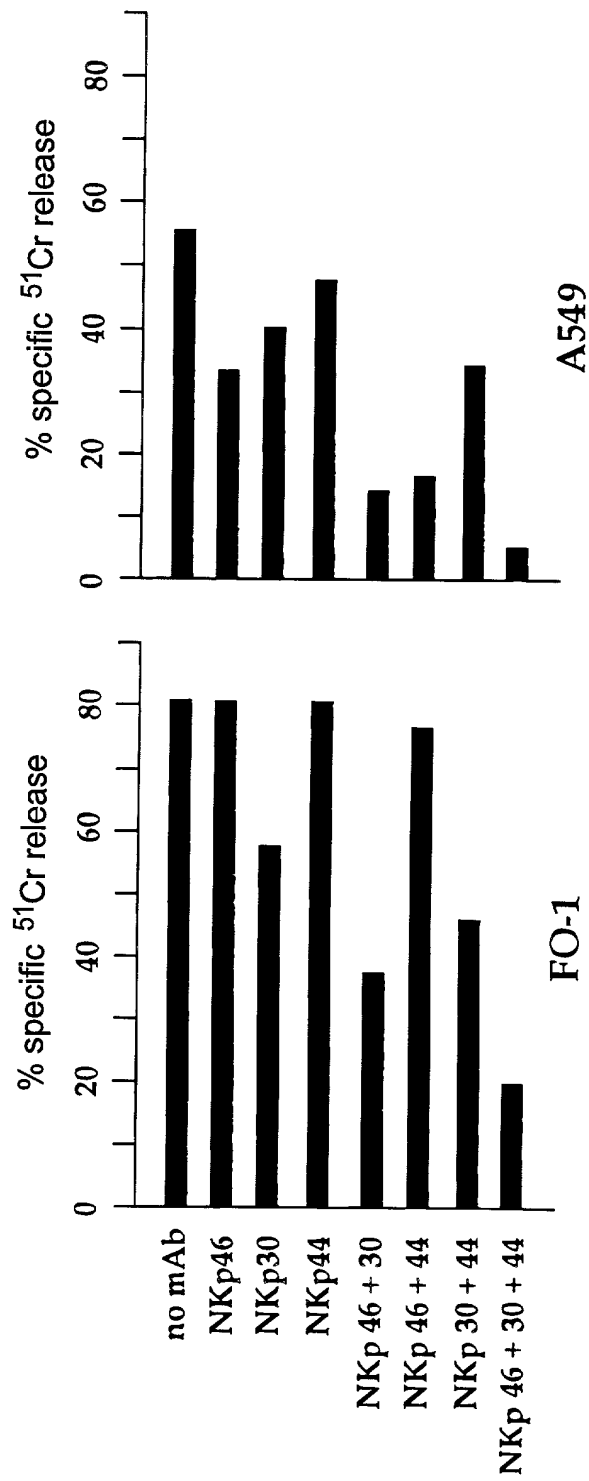


FIGURE 6B

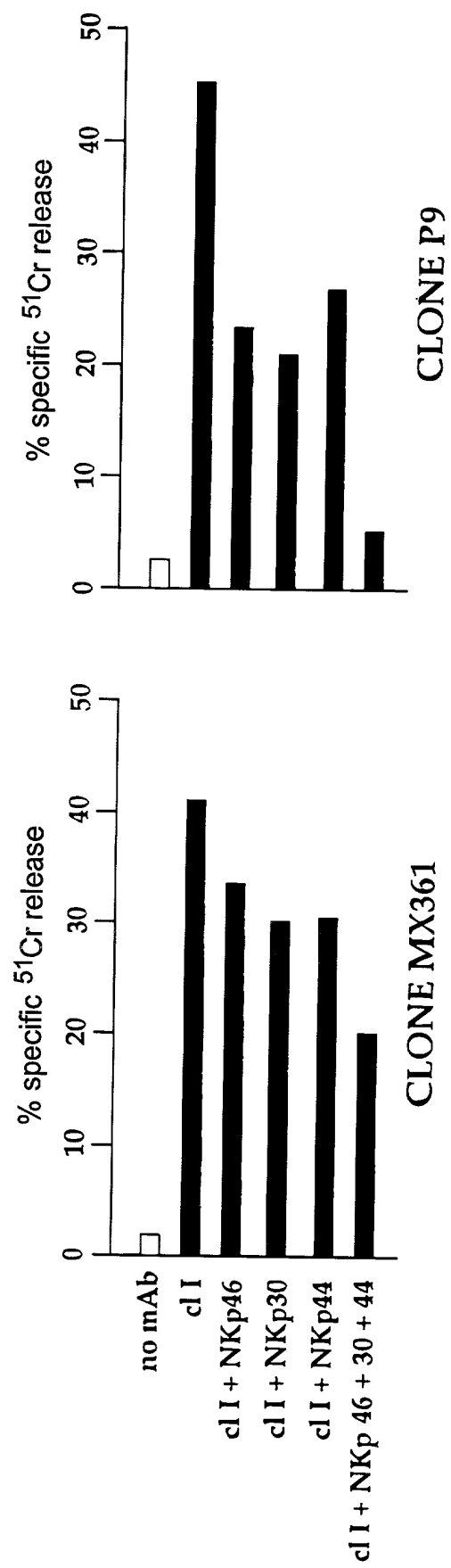


FIGURE 7A

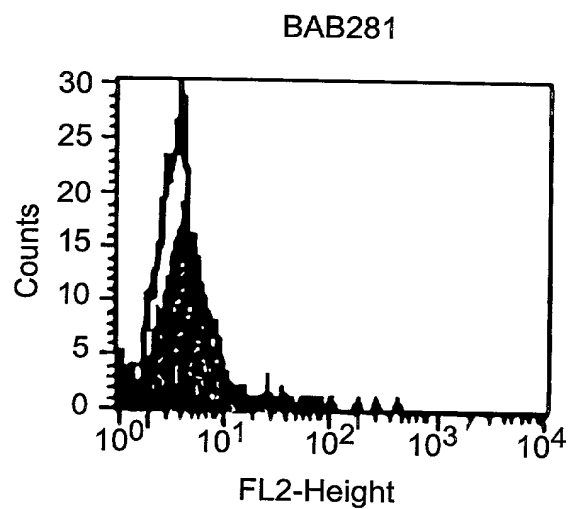
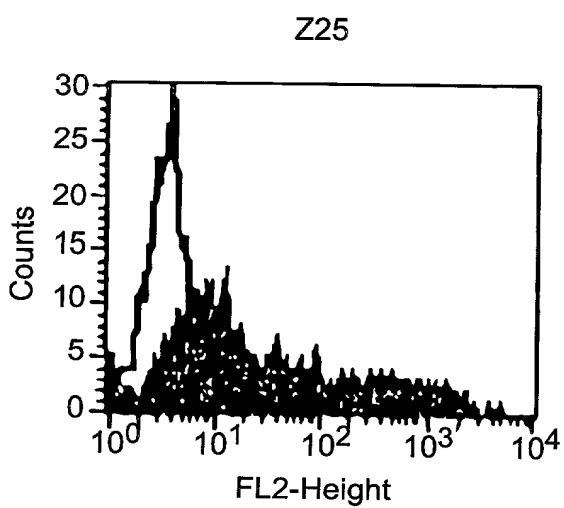
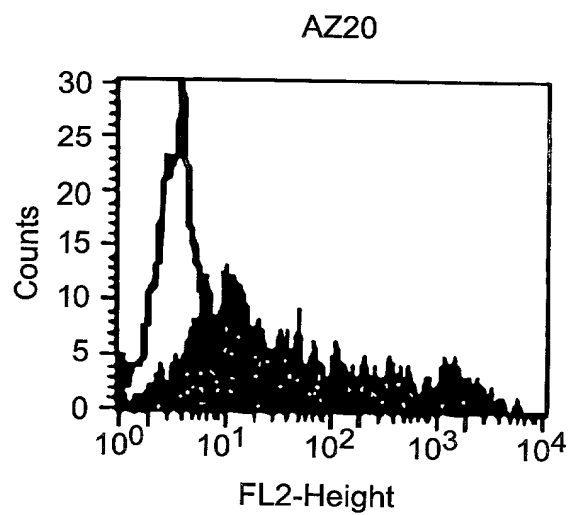
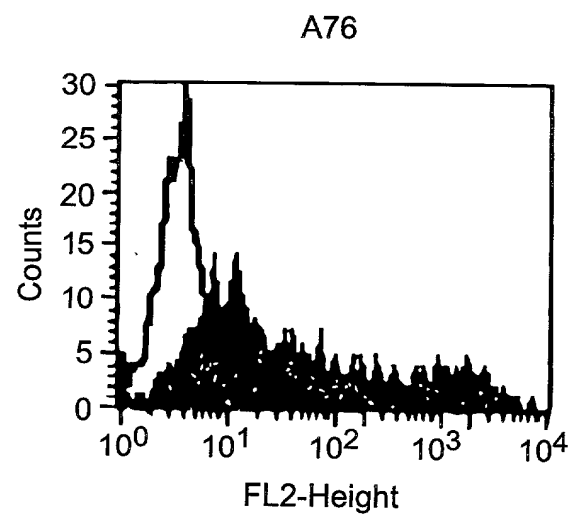


FIGURE 7B

SEQUENCE N° 2

mawmlllili	mvhpgscalW	VSQPPEIRTL	EGSSAFLPOS	<u>FN</u> ASQRLAI	50
GSVTWFRDEV	VPGKEVRNGT	PEFRGR LAPL	ASSRFLHDHQ	AELHIRDVRG	100
HDASIYVQRV	EVLGLGVGTG	<u>NG</u> TRLVVEKE	HPQLGAGTVL	<u>LLRAGFYAVS</u>	150
<u>FLS</u> VAVGSTV	YYQ GKCHCHM	GTHCHSSDGP	RGVIPEPRCP		190

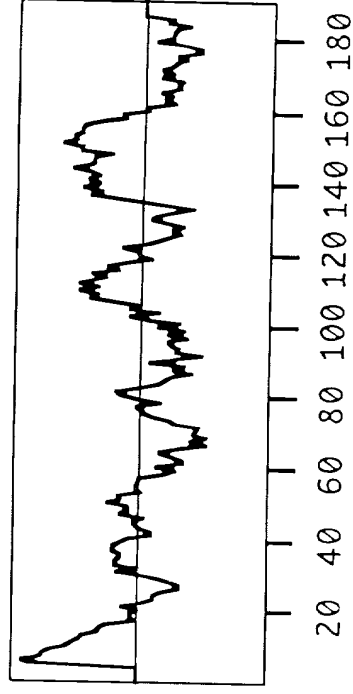


FIGURE 7C

SEQ ID N° 1

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1  ccttcctcct ccaccagac ctactgctc agacccccc cgcaactgg gacatcttcc
61  gacatggcct ggatgcctgt gctcaacttg atcatggctc atccaggac ctgctctctc
121  tgggtgtccc agcccccctga gattcgtacc ctggaaggac cctctgccc cctgcccgc
181  tccctcaatg ccagccaagg gagactggcc attggctccg caactggctt ccgagatgag
241  gtggttccag ggaaggaggt gaggaatgga acccagagt tcaggggccg cctggcccca
301  ctctgctctc ccgcttctcc ccctgacctc caggctggag tgcacatccg ggaactgaga
361  ggccatgacg ccagcatcta cgtgtgcaga gtggaggttc tgggcccctg cgtcgggaca
421  gggaaatggg ctgggttgtt ggctggagaaa gaacatcccc agctaggggc tggtaacagt
481  ctccctccctc gggctggact ctatgctgtc agcttctctc ctgtggccgc ggcagcacc
541  gctatattac agggcaaatg cactgtcac atgggaacac actgccactt ccaagatggg
601  ccccgaggcg tgaattccaga gccagatgt ccttagtctc cccaaaaga ccccaacaaa
661  tctgcccac cact

```

FIGURE 8A

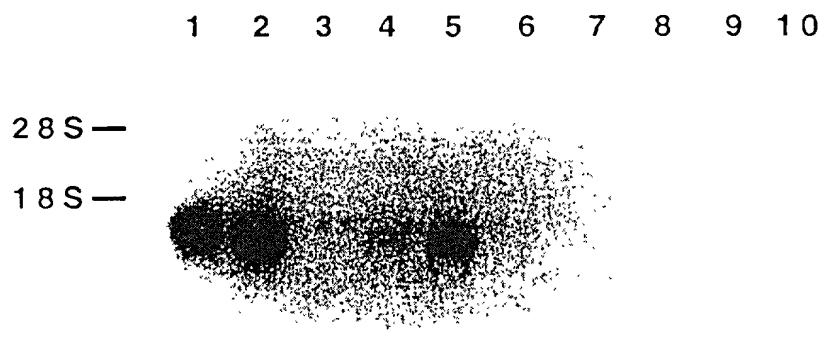


FIGURE 8B

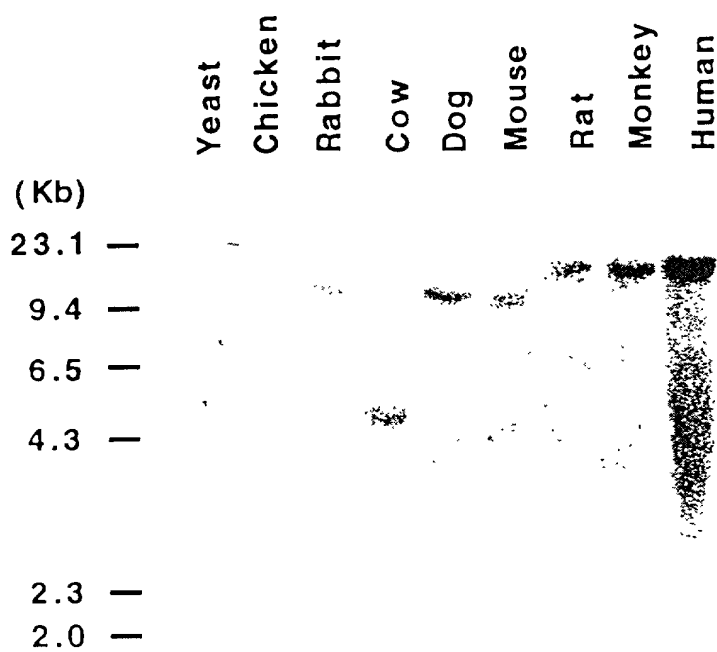
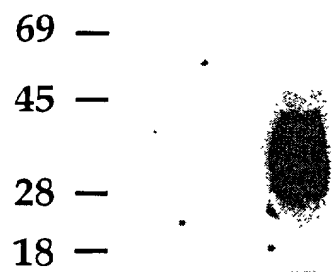


FIGURE 9A

Cells Daudi NK



Blot

I

[illegible]